

PEPTIDESTRUCTURE of: human\_prol.pap ck: 2321, 1 to 458

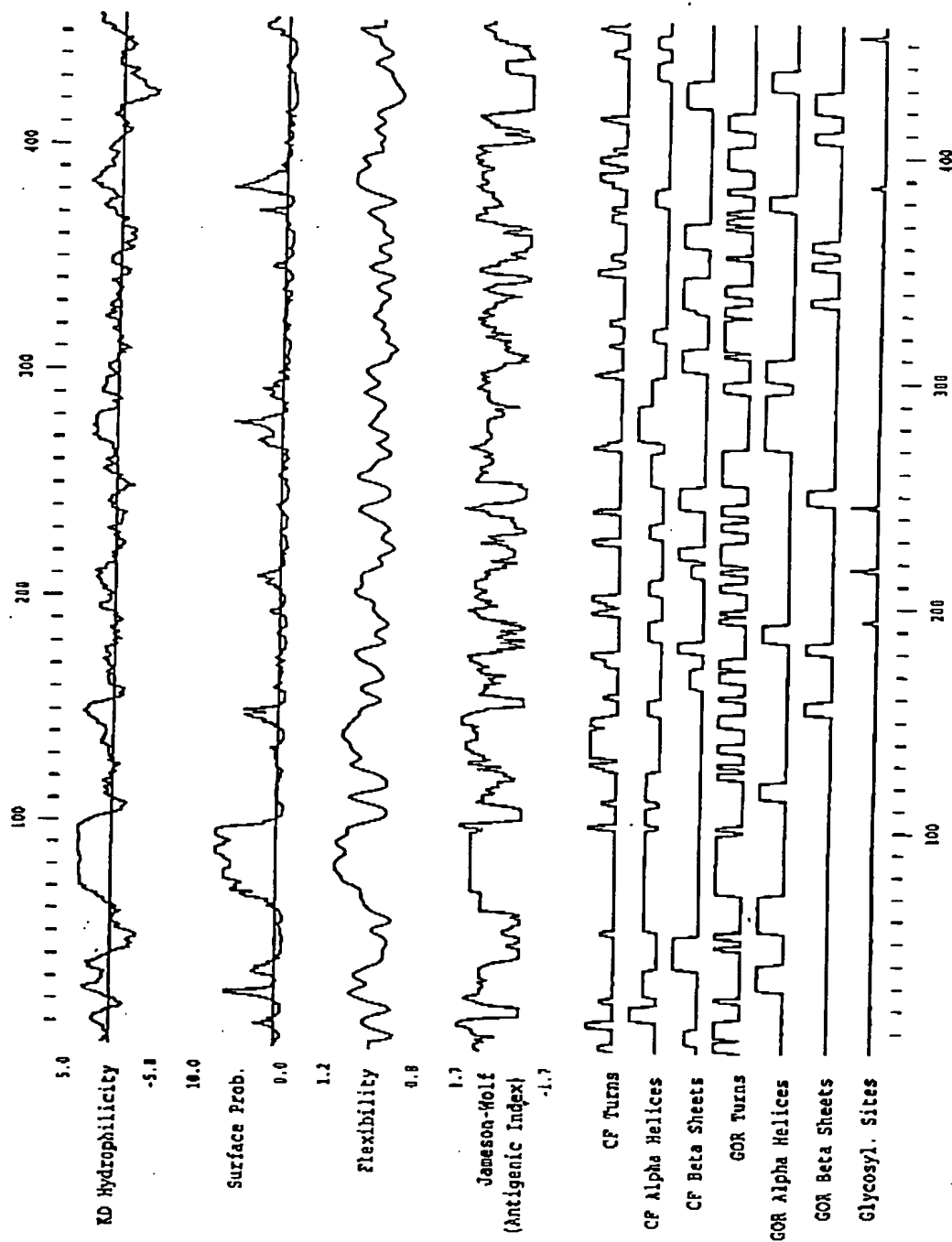
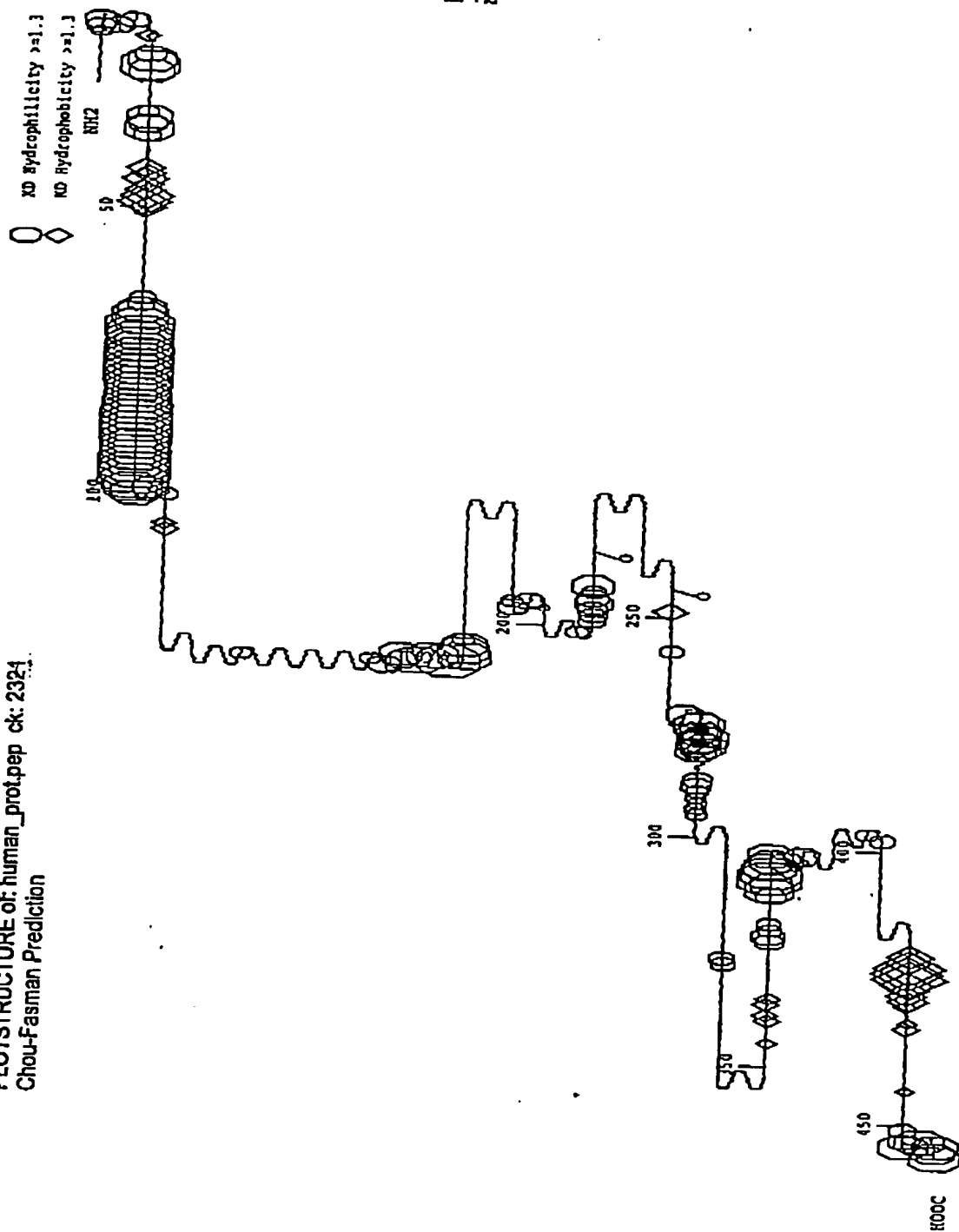


FIGURE 1

662TQT"48E9T460

PLOTSTRUCTURE of: human\_prot.pep ck: 2324  
Chou-Fasman Prediction



2/12

FIGURE 2

662TQT"48E9T460

PLOTSTRUCTURE of: human\_prot.pep ck: 2321.  
Garnier-Osguthorpe-Robson Prediction

○ Hydrophilicity >1.3  
○ Hydrophobicity >1.3

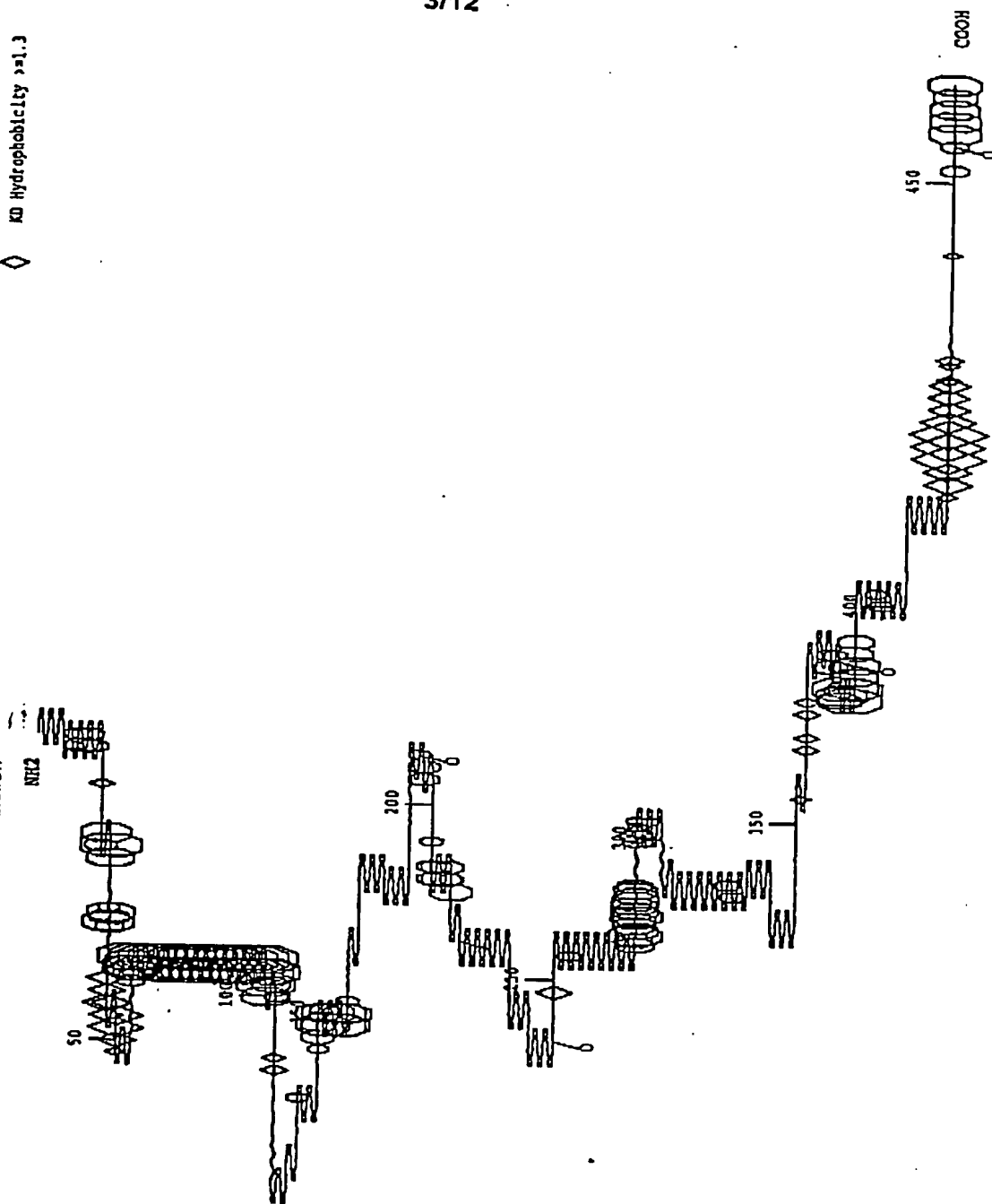


FIGURE 3

PEPTIDESTRUCTURE of: g713estmouse.0.12\_tr13.pap ck: 1880, 1 to 467

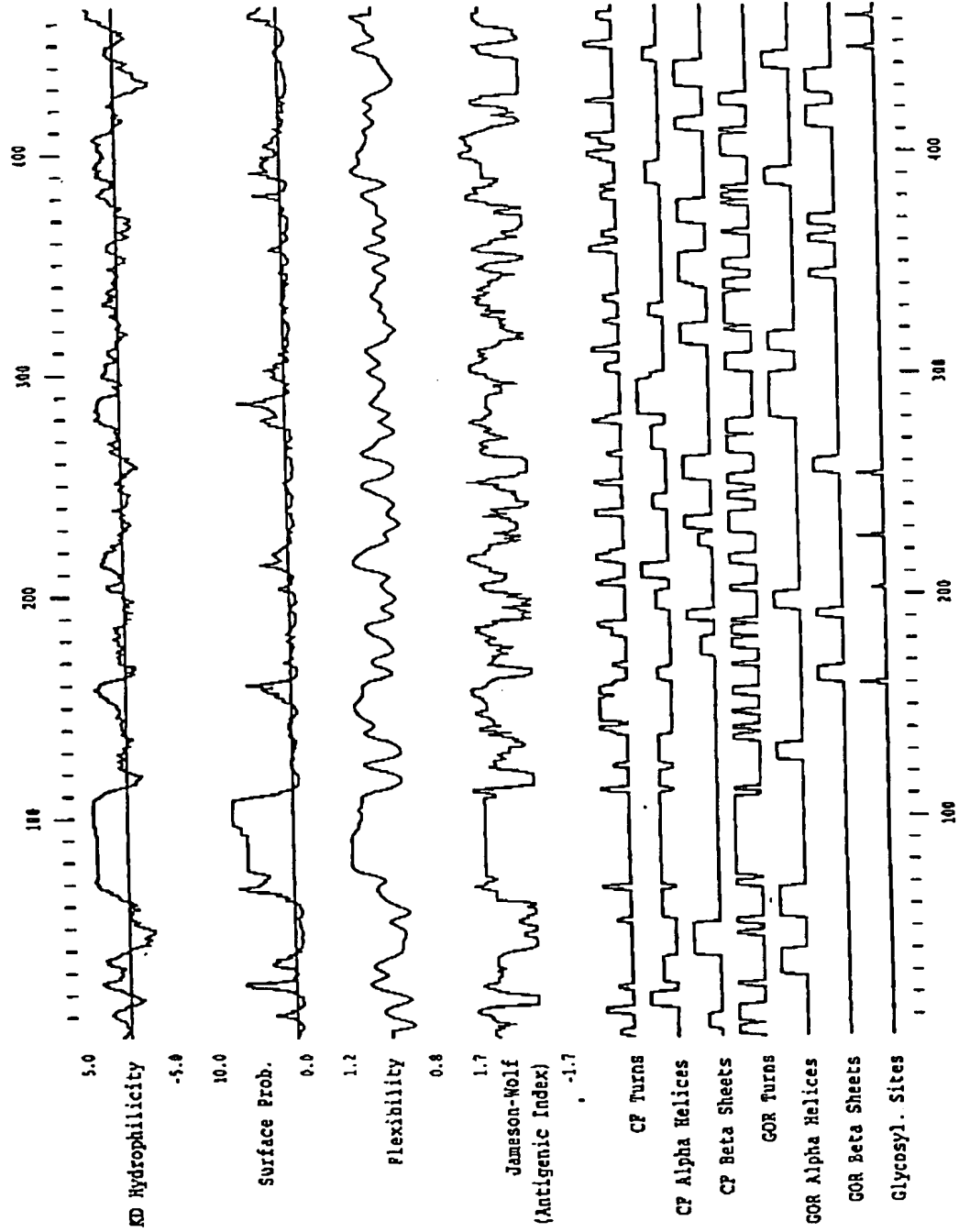


FIGURE 4

662707 "HEESTH60

PLOTSTRUCTURE of: g713estmouse.0.12\_tr13.pep ck: 1880  
Chou-Fasman Prediction

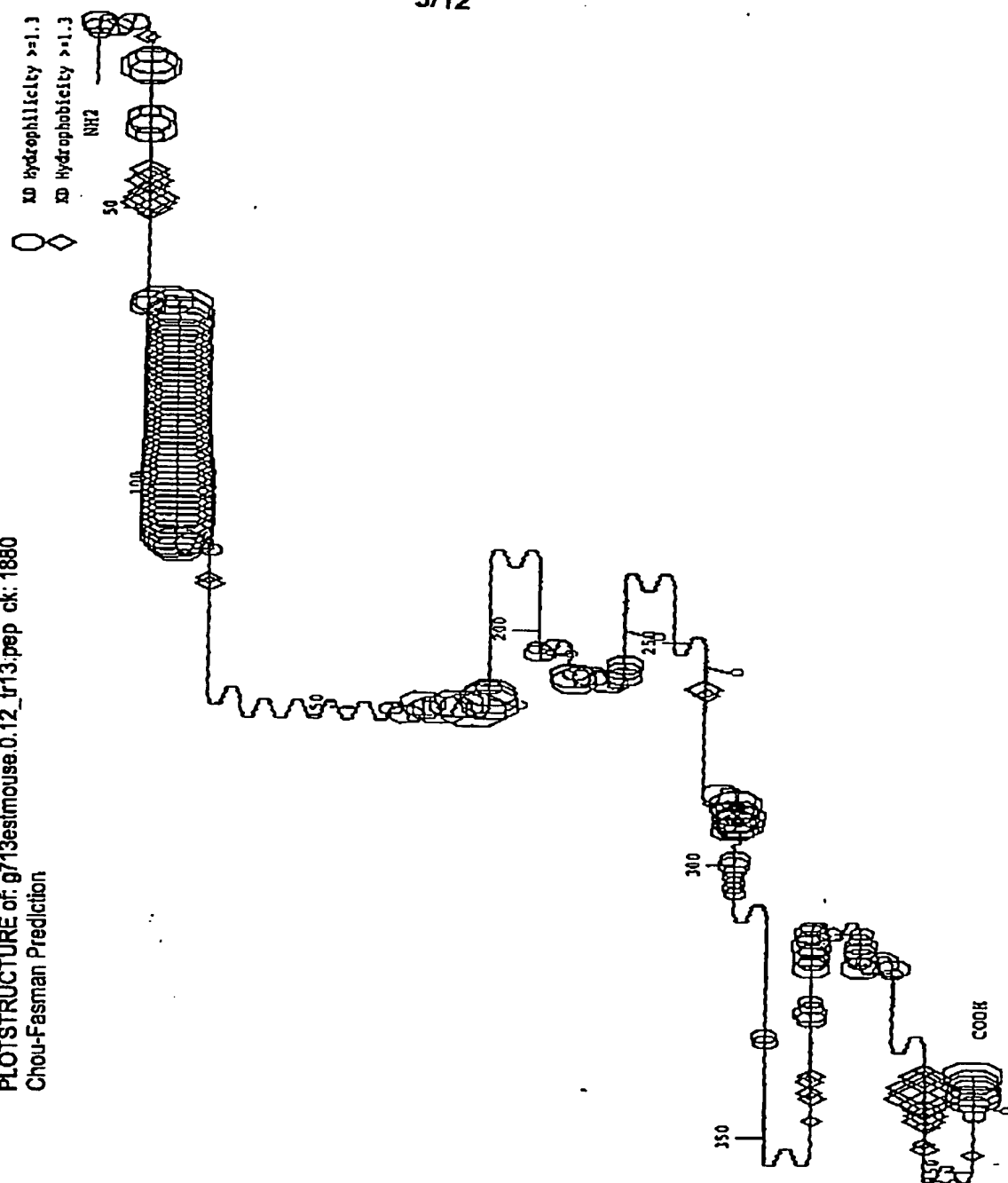


FIGURE 5

662TQT"48E9T460

PLOTSTRUCTURE of: g713estmouse.0.12\_tr13.pép ck: 1880  
Garnier-Osguthorpe-Robson Prediction

⊗ XD Hydrophilicity >=1.3  
⊗ XD Hydrophobicity >=1.3

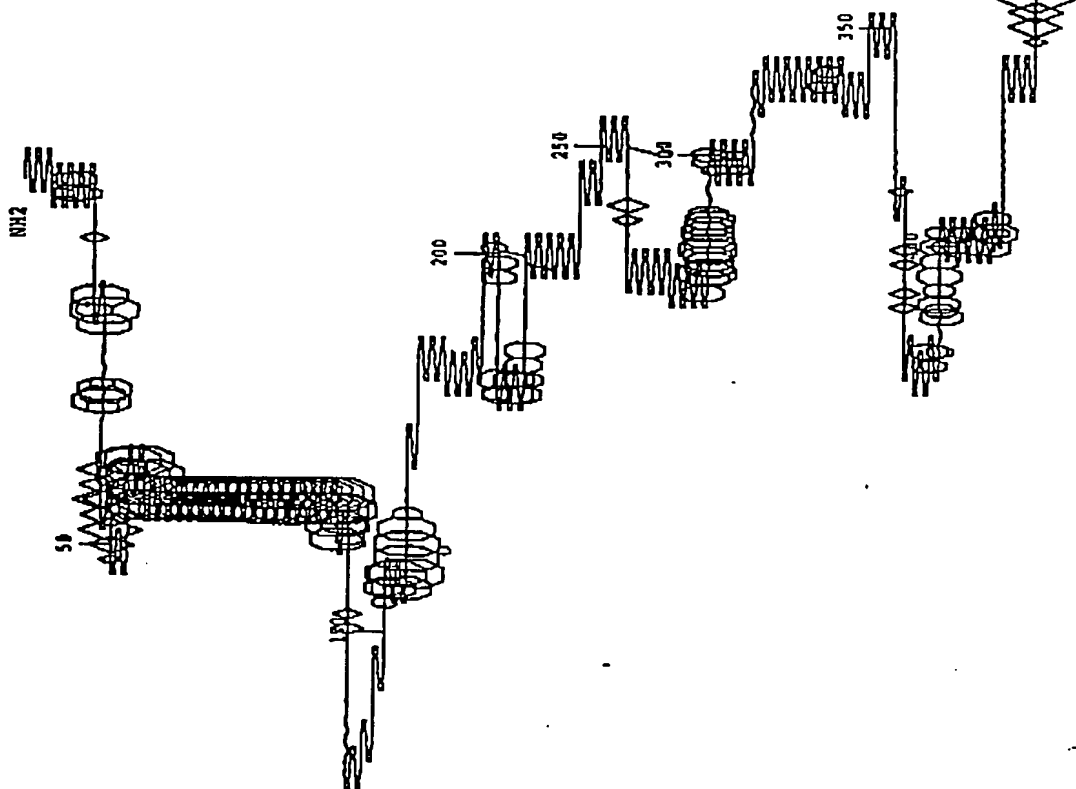


FIGURE 6

09416384.101299

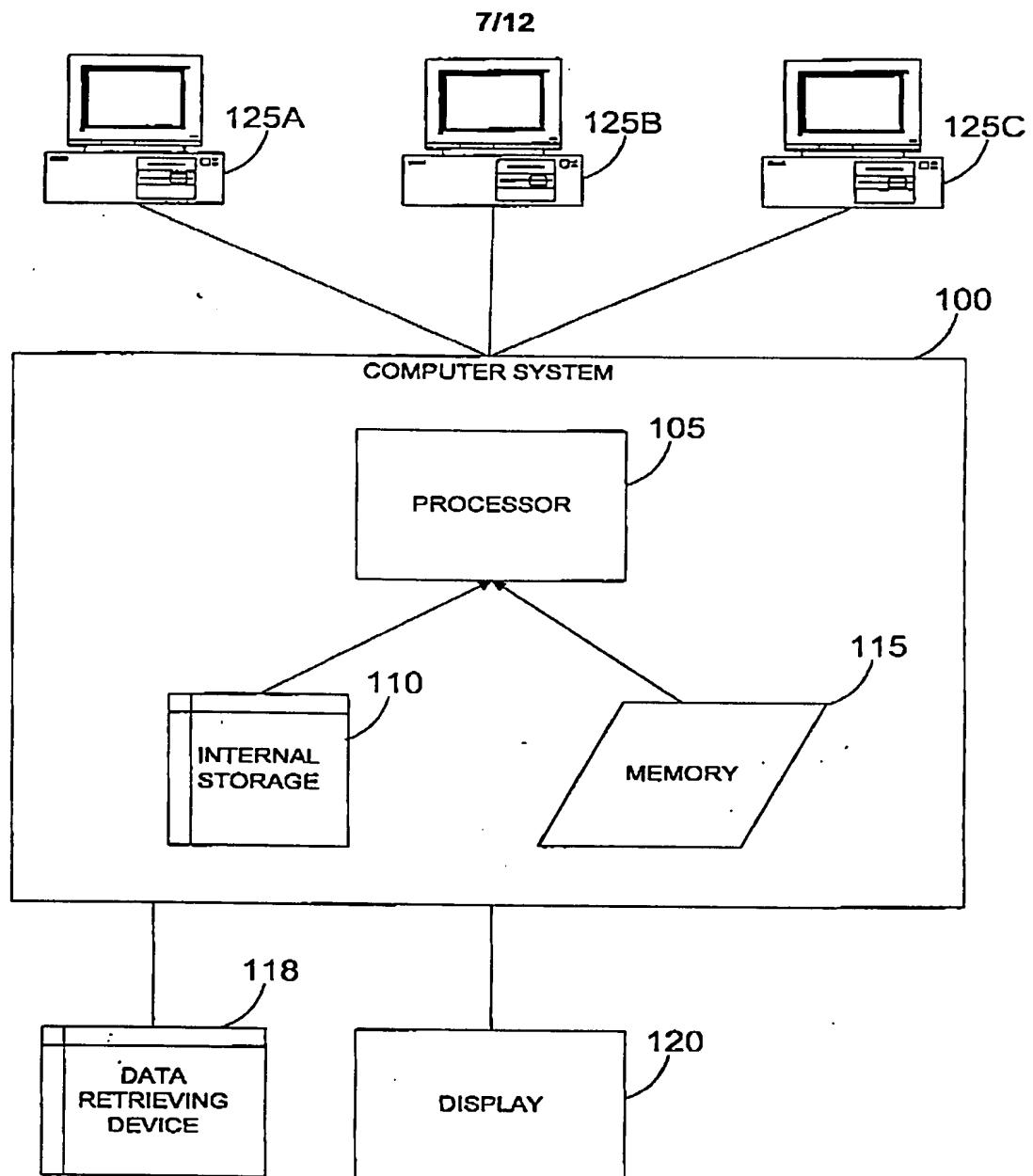


FIGURE 7

09416384.101299

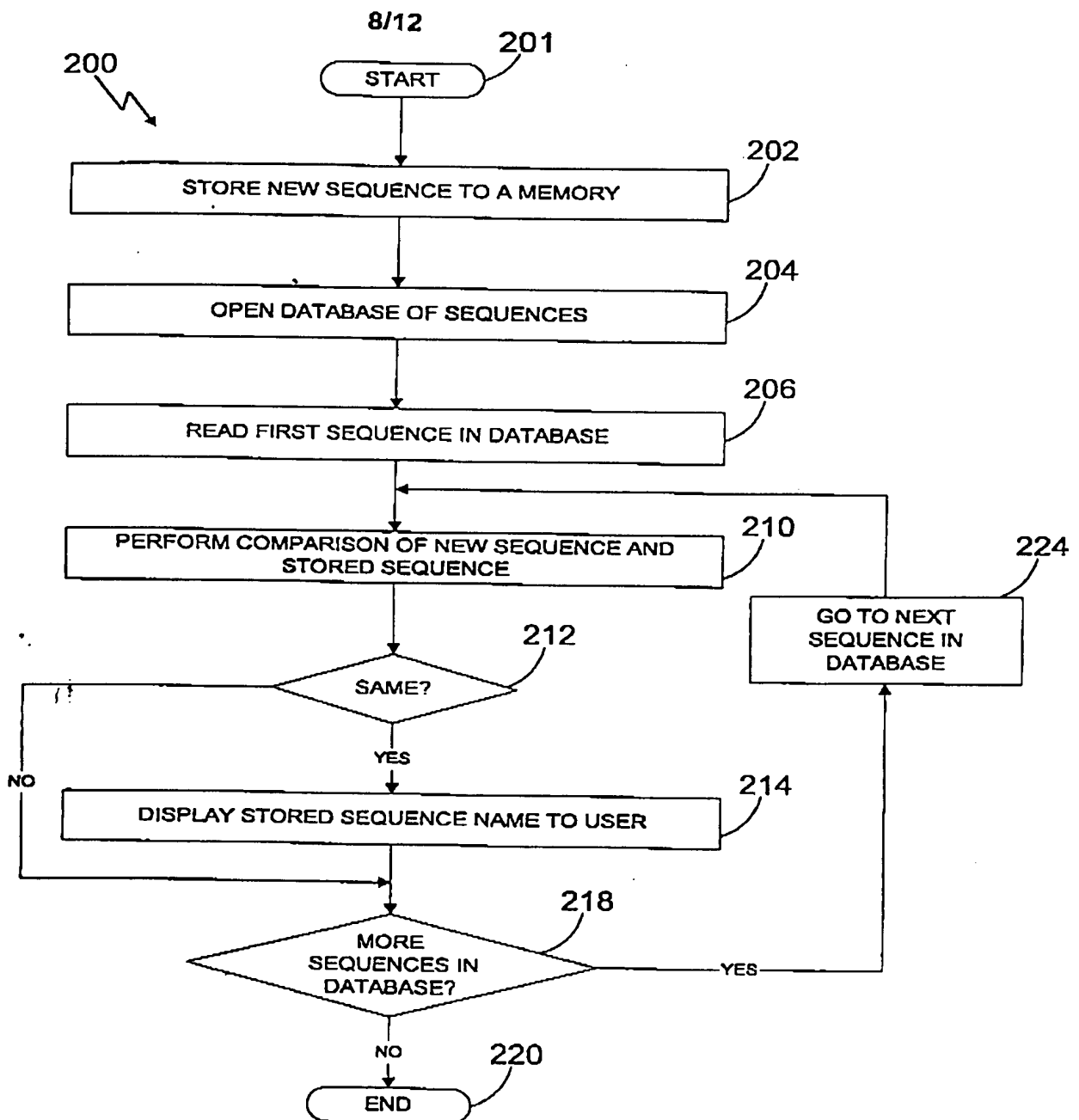


FIGURE 8



00415384-101299

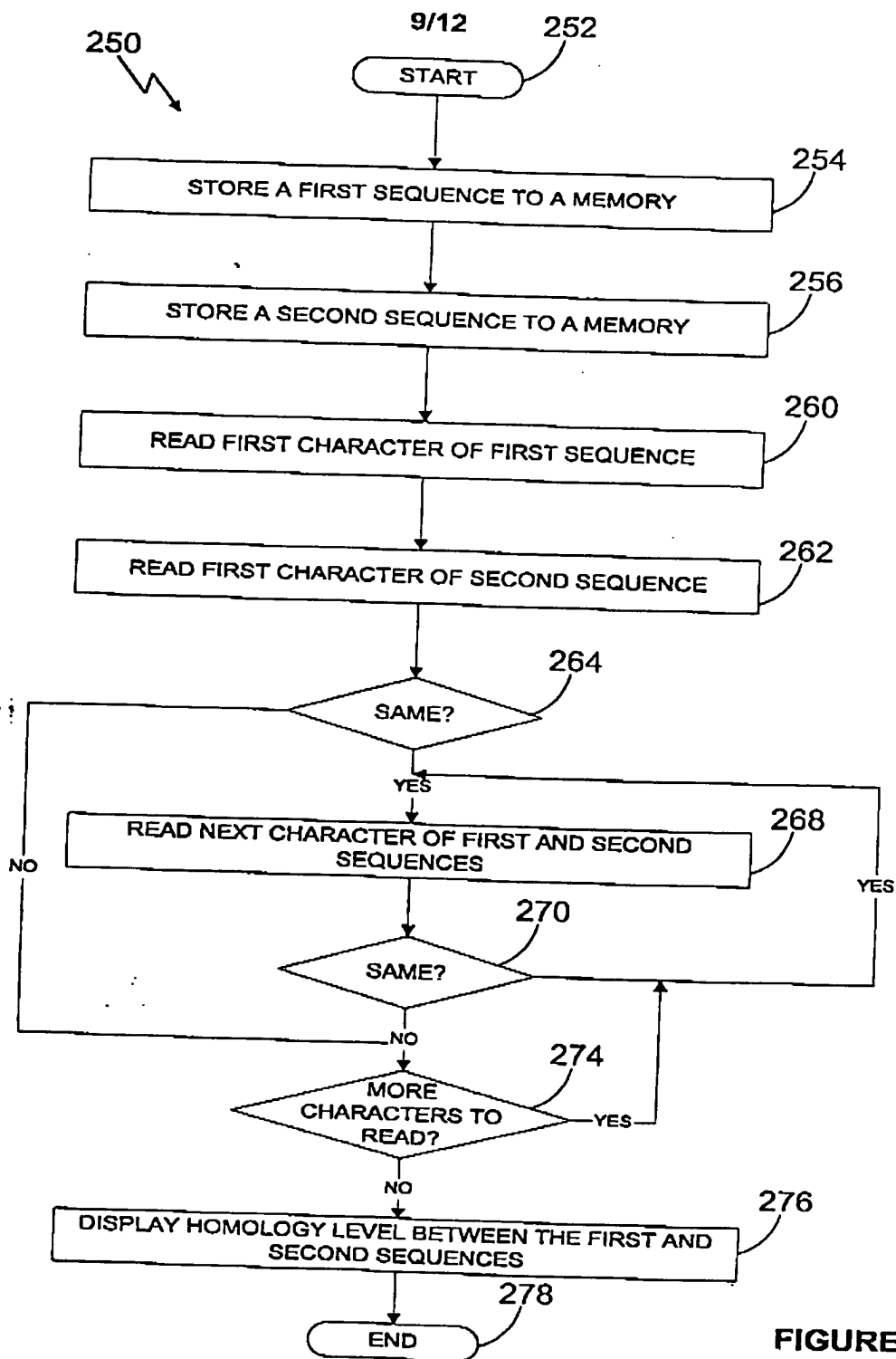


FIGURE 9

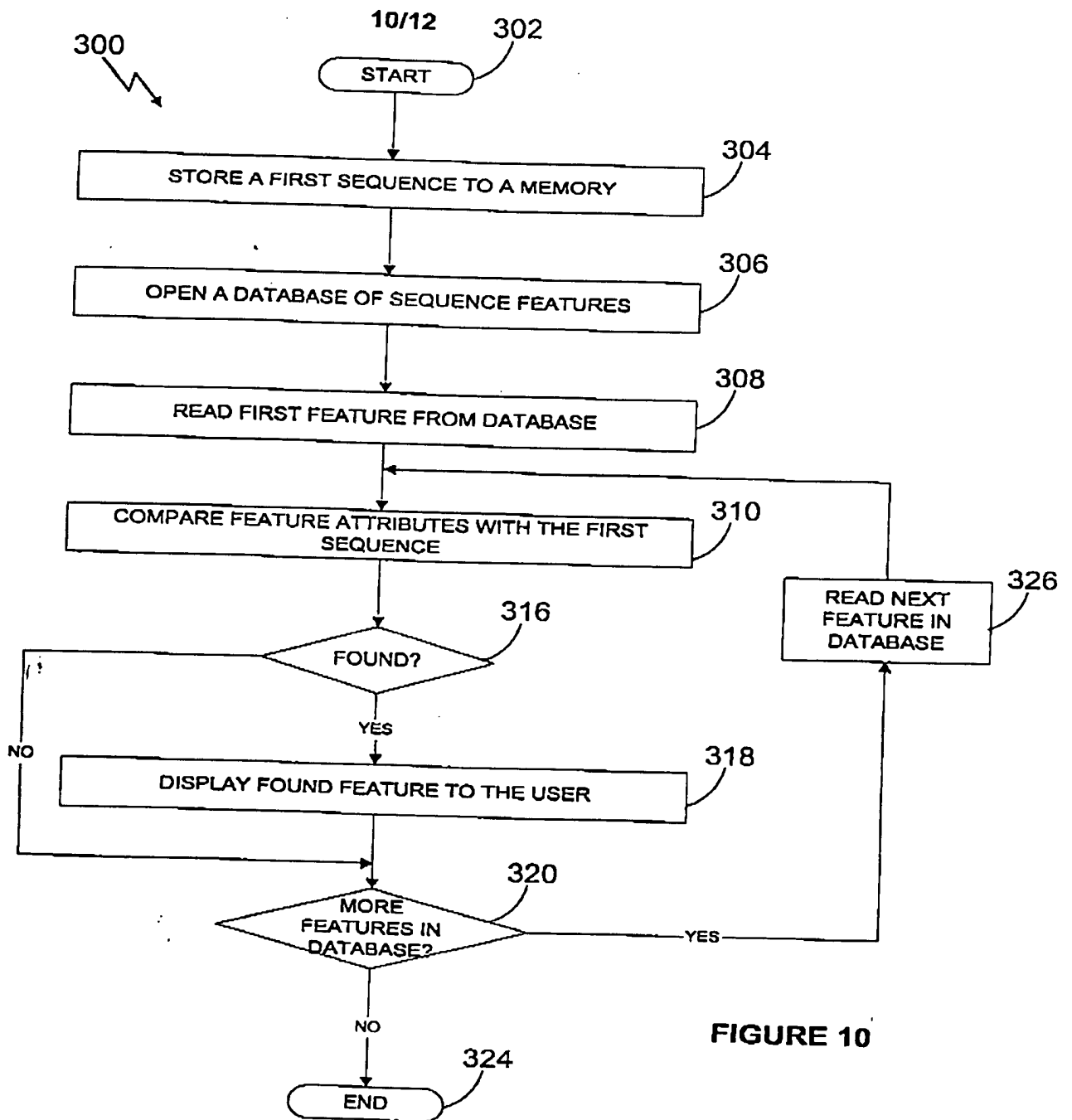


FIGURE 10

FIGURE 11

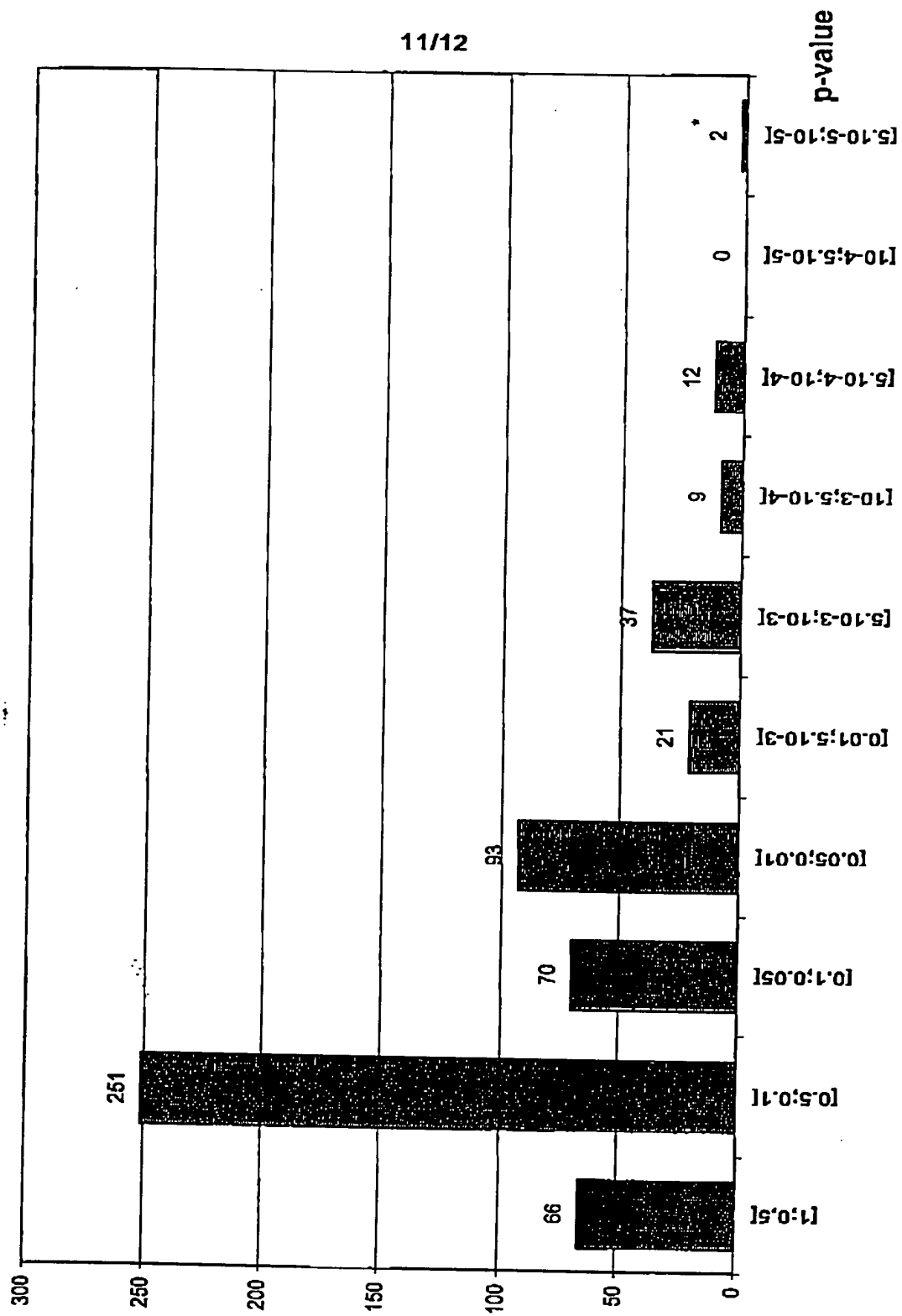


FIGURE 12

